

SEQUENCE LISTING

<110> VLAAMS INTERUNIVERSITAIR INSTITUUT VOOR BIOTECHNOL

<120> NUCLEIC ACID BINDING OF MULTI-ZINC FINGER TRANSCRIPTION FACTORS

<130> JAR/SIP/V042

<140> PCT/EP00/05582

<141> 2000-06-09

<150> 99202068.5

<151> 1999-06-25

<160> 50

<170> PatentIn Ver. 2.1

<210> 1

<211> 11

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: part of bait
for screening

<220>

<221> misc_feature

<222> (6)

<223> n is a spacer sequence of at least 8 base pairs

<400> 1

cacctncacc t

11

<210> 2

<211> 11

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: part of bait
for screening

<220>

<221> misc_feature

<222> (6)

<223> n is a spacer sequence of at least 8 base pairs

<400> 2

cacctnaggt g

11

<210> 3

<211> 11

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: part of bait for screening

<220>

<221> misc_feature

<222> (6)

<223> n is a spacer sequence of at least 8 base pairs

<400> 3

aggtgncacc t

11

<210> 4

<211> 11

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: part of bait
for screening

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<222> (6)

<223> n is a spacer sequence of at least 8 base pairs

<400> 4
aggtgnaggt g

11

<210> 5
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<212> DNA
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<223> Description of Artificial Sequence: bipartite element

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<400> 5
cacctncacc tg

12

<210> 6
<211> 25
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: complex
consensus sequence

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<222> (16)
<223> n is a spacer sequence of at the most 28 base pairs

<400> 6
gacaagataa gataanctca tcttc

25

<210> 7
<211> 30
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<223> Description of Artificial Sequence: primer SIP1\NZF3Mut

<400> 7

ccacctgaaa gaatccctga gaattcacag

30

<210> 8

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: primer SIP1
NZF4Mut

<400> 8

gggtcctaca gttcatctat cagcagcaag

30

<210> 9

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: primer SIP1 CZF2Mut

<400> 9

caccacctta tcgagtcctc gaggctgcac

30

<210> 10

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: primer SIP1
CZF3Mut

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tcctactcgc agtccatgaa tcacaggtac

30

<210> 11
<211> 50
<212> DNA
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<223> Description of Artificial Sequence: probe Xbra-WT

<400> 11
atccaggcca cctaaaatat agaatgataa agtgaccagg tgtcagttct 50

<210> 12
<211> 50
<212> DNA
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<400> 12
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<210> 13
<211> 23
<212> DNA
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<223> Description of Artificial Sequence: probe Xbra-E

<400> 13
taaagtgacc aggtgtcagt tct 23

<210> 14
<211> 27
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<223> Description of Artificial Sequence: probe Xbra-F

<400> 14

atccaggcca cctaaaatat agaatga

27

<210> 15

<211> 50

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Rdm + Xbra-E

<400> 15

caatttagag tactgtgtac ttggggagtaa agtgaccagg tgtcagttct

50

<210> 16

<211> 53

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: probe Xbra-F + AREB6

<400> 16

atccaggcca cctaaaatat agaatgaggc tcagacaggt gtagaattcg gcg

53

<210> 17

<211> 53

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Rdm + AREB6

<400> 17

caatttagag tactgtgtac ttggggagggc tcagacaggt gtagaattcg gcg

53

<210> 18

<211> 50

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: probe Xbra-J

<400> 18

gcacaggcca cctaaaatat agaatgataa agtgaccagg tgtcagttct 50

<210> 19

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Xbra-K

<400> 19

atcactgcca cctaaaatat agaatgataa agtgaccagg tgtcagttct 50

<210> 20

<211> 50

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: probe Xbra-L

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<210> 21

<211> 50

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: probe Xbra-M

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atccaggccc aataaaatat agaatgataa agtgaccagg tgtcagttct 50

<210> 22

<211> 50

<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: probe Xbra-N

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atccaggcca ccgccaatat agaatgataa agtgaccagg tgtcagttct 50

<210> 23
<211> 50
<212> DNA
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<223> Description of Artificial Sequence: probe Xbra-O

<400> 23
atccaggcca cctaaccgat agaatgataa agtgaccagg tgtcagttct 50

<210> 24
<211> 50
<212> DNA
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<223> Description of Artificial Sequence: probe Xbra-P

<400> 24
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<210> 25
<211> 50
<212> DNA
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<223> Description of Artificial Sequence: probe Xbra-Q

<400> 25
atccaggcca cctaaaatat atcctgataa agtgaccagg tgtcagttct 50

<210> 26

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Xbra-R

<400> 26

atccaggcca cctaaaatat agaagtctaa agtgaccagg tgtcagttct 50

<210> 27

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Xbra-S

<400> 27

atccaggcca tctaaaatat agaatgataa agtgaccagg tgtcagttct 50

<210> 28

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Xbra-Z

<400> 28

atccaggcca cctaaaatat agaatgataa agtgactagg tgtcagttct 50

<210> 29

<211> 47

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Xbra-B

<400> 29
atccaggcca cctatataga atgataaagt gaccaggtgt cagttct 47

<210> 30
<211> 47
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: probe Xbra-C

<400> 30
atccaggcca cctaaaatat agaatgatgt gaccaggtgt cagttct 47

<210> 31
<211> 40
<212> DNA
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<220>
<223> Description of Artificial Sequence: probe Xbra-U

<400> 31
atccaggcca cctaaaatat agtgaccagg tgtcagttct 40

<210> 32
<211> 46
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: probe Xbra-EE

<400> 32
taaagtgacc aggtgtcagt tcttaaagtg accaggtgtc agttct 46

<210> 33
<211> 46
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Xbra-ErE

<400> 33

agaactgaca cctgggtcact ttataaagtg accaggtgtc agttct

46

<210> 34

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Xbra-FrF

<400> 34

atccaggcca cctaaaatat agaatattct atatttttagg tggcctggat

50

<210> 35

<211> 50

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: probe Xbra-V

<400> 35

atccaggcag gtgtaaatat agaatgataa agtgaccac ctacagttct

50

<210> 36

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Xbra-W

<400> 36

atccaggcag gtgtaaatat agaatgataa agtgaccagg tgtcagttct

50

<210> 37

<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: probe alfa4I-WT (alfa-4-integrin)

<400> 37
gcagggcaca cctggattgc attagaatga gactcactac ccagttcagg tgtgttgcgt 60

<210> 38
<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: probe alfa4I-A (alfa-4-integrin)

<400> 38
gcagggcaca cctggattgc attagaatga gactcactac ccagttcaga tgtgttgcgt 60

<210> 39
<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: probe alfa4I-B
(alfa-4-integrin)

<400> 39
gcagggcaca tctggattgc attagaatga gactcactac ccagttcagg tgtgttgcgt 60

<210> 40
<211> 70
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: probe Ecad-WT

<400> 40

tggccggcag gtgaaccctc agccaatcag cggtagggg ggcggtgctc cggggctcac 60
ctggctgcag 70

<210> 41

<211> 70

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Ecad-A

<400> 41

tggccggcag gtgaaccctc agccaatcag cggtagggg ggcggtgctc cggggctcat 60
ctggctgcag 70

<210> 42

<211> 70

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: probe Ecad-B

<400> 42

tggccggcag atgaaccctc agccaatcag cggtagggg ggcggtgctc cggggctcac 60
ctggctgcag 70

<210> 43

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR-primer

<400> 43

acaaaagaac tcagccaagt g

21

<210> 44

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR-primer

<400> 44

ccgcaagctc acaggtgc

18

<210> 45

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: forward primer E-box1

<400> 45

gctgtggccg gcagatgaac cctcag

26

<210> 46

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: reverse primer E-box1

<400> 46

ctgagggttc atctgccggc cacagc

26

<210> 47

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: forward primer
E-box3

<400> 47

gctccgggct catctggctg cagc

24

<210> 48

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: reverse primer E-box3

<400> 48

gctgcagcca gatgagcccc ggagc

25

<210> 49

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: degenerated primer

<400> 49

cttccagcag ccctacgayc argcnca

27

<210> 50

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: degenerated primer

<400> 50

gggtgtggga ccggatrtgc atyttnat

28
